

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/562,627A
Source: TFWP
Date Processed by STIC: 07/10/2006

ENTERED



IFWP

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/562,627A

DATE: 07/10/2006

TIME: 10:45:45

Input Set : F:\Substitute CRF Seq Listing 6-05-2006.txt

Output Set: N:\CRF4\07102006\J562627A.raw

3 <110> APPLICANT: CHOE, Mu-Hyeon
 4 CHOI, Seong-Hyeok
 5 LEE, Yong-Chan
 6 KWON, Hye-Won
 7 WON, Jae-Seon
 8 YU, Mi-Hyun
 9 SONG, Jeong-Hwa
 10 KIM, Yong-Jae
 12 <120> TITLE OF INVENTION: The Dimer of Chimeric Recombinant Binding
 13 Domain-Functional Group Fusion formed via
 14 Disulfide-bond-bridge and The Process
 15 For Producing The Same
 17 <130> FILE REFERENCE: 428.1060
 19 <140> CURRENT APPLICATION NUMBER: US 10/562,627A
 20 <141> CURRENT FILING DATE: 2005-12-22
 22 <150> PRIOR APPLICATION NUMBER: PCT/KR2004/001595
 23 <151> PRIOR FILING DATE: 2004-06-30
 25 <150> PRIOR APPLICATION NUMBER: KR2003-0043599
 26 <151> PRIOR FILING DATE: 2003-06-30
 28 <160> NUMBER OF SEQ ID NOS: 12
 30 <170> SOFTWARE: KopatentIn 1.71
 32 <210> SEQ ID NO: 1
 33 <211> LENGTH: 1749
 34 <212> TYPE: DNA
 35 <213> ORGANISM: Artificial Sequence
 W--> 36 <220> FEATURE:
 37 <223> OTHER INFORMATION: pMC74 plasmid coding sequence
 39 <400> SEQUENCE: 1

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44	actccagaga	agaggctgga	gtgggtcgca	tacatttagta	atgatgatag	ttccggccgt	180
46	tattcagaca	ctgtaaaaggg	ccgggttacc	atctccagag	acaatgcgcag	gaacaccctc	240
48	tacctgcaaa	tgagccgtct	gaagtctgag	gacacagcca	tatattctgt	tgcaagagga	300
50	ctggcctggg	gagctctggtt	tgcttactgg	ggccaaaggga	ctctggtcac	tgtctctgca	360
52	gccaaaacga	caccccccattc	tgtctatcca	ctggccctgt	gatctgcgtc	ccaaactaac	420
54	tccatggtga	ccctggatg	cctggtcaag	ggcttattcc	ctgagccagt	gacagtgacc	480
56	tggaactctg	gatccctgtc	cagcgggtgt	cacacattcc	cagctgtcct	gcagtctgac	540
58	ctctacactc	tgaggcagctc	agtgactgtc	ccctccagca	cctggcccgag	cgagaccgtc	600
60	acctgcaacg	ttgcccaccc	ggccagcagc	accaagggtgg	acaagaaaaat	tgtgcccagg	660
62	gattgtggta	gtaaggcttag	cataagtaca	aaagcttccg	gagggtcccg	ggggggcagc	720
64	ctggccgcgc	tgaccgcgc	ccaggcttgc	cacctggccgc	tggagacttt	cacccgtcat	780
66	cgccagccgc	gccccgtggga	acaactggag	cagtgcggct	atccggtgca	gccccgtggc	840
68	gcccttacc	tggccgcgc	gctgtcggtt	aaccaggatcg	accaggatcg	ccgcaacgc	900

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74	ggcaaacacg	aggccggcgc	ggccaacggc	ccggcggaca	gcccgcacgc	cctgctggag	1080
76	cgcaactatc	ccactggcgc	ggagttcctc	ggcgaacggc	gcccacgtcag	cttcagcacc	1140
78	cgccggcacgc	agaactggac	ggtggagcgg	ctgctccagg	cgaccggcca	actggaggag	1200
80	cgccggctatg	tgttcgtcgg	ctaccacggc	accttctcg	aagcggcgcga	aagcatcgtc	1260
82	ttcggcgggg	tgcgcgcgcg	cagccaggac	ctcgacgcga	tctggcgcgg	tttctatatc	1320
84	gccggcgatc	cgccgcgtggc	ctacggctac	gcccaggacc	aggaacccga	cgcacgcggc	1380
86	cgatccgca	acgggtccct	gctgcggggtc	tatgtgccgc	gctcgagcct	gccggcgttc	1440
88	taccgcacca	gcctgaccct	ggccgcgcgg	gaggcggcgg	gcccgggtcga	acggctgatc	1500
90	ggccatccgc	tgcgcgtcgc	cctggacgc	atcaccggcc	ccgaggagga	aggcgggcgc	1560
92	ctggagacca	ttctcggtc	gcccgcggcc	gagcgcaccc	tggtgattcc	ctcggcgatc	1620
94	ccaccggacc	cgcgcaacgt	cgccggcgcac	ctcgaccctgt	ccagcatccc	cgacaaggaa	1680
96	caggcgatca	gcccctgcc	ggactacgc	agccagcccg	gcaaaccggc	gcccggaggac	1740
98	ctgaagtaa						1749

101 <210> SEQ ID NO: 2

102 <211> LENGTH: 1764

103 <212> TYPE: DNA

104 <213> ORGANISM: Artificial Sequence

W--> 105 <220> FEATURE:

106 <223> OTHER INFORMATION: pMH21 plasmid coding sequence

108 <400> SEQUENCE: 2

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113	actccagaga	agaggctgg	gtgggtcgca	tacatttagt	atgtatgtat	ttccgcgcgt	180
115	tattcagaca	ctgttaaagg	ccgggttcc	atctccagag	acaatgc	gaacaccctc	240
117	tacctgcaaa	tgagccgtc	gaagtctgag	gacacagcc	tatattcctg	tgcaagagga	300
119	ctggcctggg	gagcctgg	tgcttactgg	ggccaaggga	ctctggtcac	tgtctctgca	360
121	gccaaaacga	caccatccatc	tgttattcca	ctggccctg	gatctgtc	ccaaactaac	420
123	tccatggta	ccctggat	cctggtaaag	ggctatttcc	ctgagccagt	gacagtgacc	480
125	tggaactctg	gatccctgtc	cagcgggtgt	cacacccctc	cagctgtc	gcaatctgac	540
127	ctctacactc	tgagcagctc	agtactgtc	ccctccagca	cctggcc	cgagaccgtc	600
129	acctgcaacg	ttgcccaccc	ggccagcagc	accaagggt	acaagaaaat	tgtcccagg	660
131	gattgtggta	gtaagcctt	cataagtaca	aaagcttctg	gtgggtggcgg	atctggaggt	720
133	cccgaggccg	gcagcctggc	cgcgtgacc	gcccaccagg	cttgcac	gcccgtggag	780
135	actttcaccc	gtcatcgca	ggcgcgcggc	tggaaacaac	tggagcgt	cggtatccg	840
137	gtcagcggc	tgggtccct	ctacctggcg	gcccgggtgt	cgtggaaacca	ggtcgaccag	900
139	gtgatccca	acgcccctgg	cagcccccgg	agccggcggc	acctggcg	agcgatccgc	960
141	gagcagccgg	agcaggcccg	tctggccctg	accctggccg	ccggccgagag	cgagcgcttc	1020
143	gtccggcagg	gcaccggca	cgacgaggcc	ggcgcggcc	acggccggc	ggacagcggc	1080
145	gacccctgc	tggagcggca	ctatccact	ggcgcggag	tcctcgcc	cgccggcgcac	1140
147	gtcagctca	gcacccggc	cacgcagaac	tggacgggt	agcggctgct	ccaggcgcac	1200
149	cgccaactgg	aggagcgcgg	ctatgtttc	gtccgttacc	acggcacctt	cctcgaaagcg	1260
151	gcccggat	gcggccggat	ccgcaacgg	gcccgttgc	aggacccctg	cgcgatctgg	1320
153	cccgacgcac	gcccggat	ccgcaacgg	gcccgttgc	gggtctatgt	gcccgttgc	1380
155	agcctggcgg	gcttctaccg	caccagcctg	accctggccg	gcccggaggc	ggcggggcgg	1440
157	gtcgaacggc	tgtcgccca	tccgttgc	ctcgccctgg	acgcccattac	cgccccggag	1500
159	gaggaaggcg	ggccctgg	gaccatttctc	ggctggccgc	tggccgagcg	caccgtgg	1560
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 165 atccccgaca aggaacaggc gatcagcgc ctgcggact acgccagcca gccccggcaaa 1740
 167 cccgcgcgc aggacactgaa gtaa 1764

170 <210> SEQ ID NO: 3

171 <211> LENGTH: 1749

172 <212> TYPE: DNA

173 <213> ORGANISM: Artificial Sequence

W--> 174 <220> FEATURE:

175 <223> OTHER INFORMATION: pCE2 plasmid coding sequence

177 <400> SEQUENCE: 3

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 182 actccagaga agaggctgga gtgggtcgca tacatttagt atgatgatag ttccggccgt 180
 184 tattcagaca ctgttaaaggg ccgggttccacc atctccagag acaatgcagg gaacaccctc 240
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 198 acctgcaacg ttgcccaccc ggccagcagg accaagggtgg acaagaaaat tgcgtccagg 660
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 228 gcccattccgc tgccgtcgcc cctggacgcc atcaccggcc cggaggagga aggcggggcgc 1560
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 232 cccaccggacc cggcgcacgc cggcggcgc acctgcacccgt ccagcatccc cgacaaaggaa 1680
 234 caggcgtatc ggcgcctgcg ggactacgcg agccagcccg gcaaaaccgcg gggcggaggac 1740
 236 ctgaagtaa 1749

239 <210> SEQ ID NO: 4

240 <211> LENGTH: 672

241 <212> TYPE: DNA

242 <213> ORGANISM: Artificial Sequence

W--> 243 <220> FEATURE:

244 <223> OTHER INFORMATION: pMC75 plasmid coding sequence

246 <400> SEQUENCE: 4

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 249 tccatcttgc cggatcttgc tgcacatccat gtttgcgttcc ttcggcgtatc 120

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 253 ttttctgggg tccccagacag gttcagtggc agtggatcag ggacagattt cacactcaag 240
 255 atcagcagag tggaggctga gatatctggg gtttattact gcttcaagg ttcacatgtt 300
 257 coattcacgt tcggctcggg gacaaagttt gaaataaaac gggctgatgc tgcaccaact 360
 259 gtatccatct tcccaccatc cagtggcag ttaacatctg gaggtgcctc agtctgtgc 420
 261 ttcttgaaca acttctaccc caaagacatc aatgtcaagt ggaagattga tggcagtgaa 480
 263 cgacaaaatg gcgtcctgaa cagttggact gatcaggaca gcaaagacag caccctacgc 540
 265 atgagcagca ccctcacgtt gaccaaggac gagttatgaac gacataacag ctatacctgt 600
 267 gaggccactc acaagacatc aacttcaccc attgtcaaga gcttcaacag gaatgagtgt 660
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272 <210> SEQ ID NO: 5

273 <211> LENGTH: 2454

274 <212> TYPE: DNA

275 <213> ORGANISM: Artificial Sequence

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277 <223> OTHER INFORMATION: pLSC52 plasmid coding sequence

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348	ctggcctacg	gctacgccc	ggaccaggaa	cccgacgcac	gcccggat	ccgcaacggt	2100
350	gccctgtgc	gggtctatgt	gcccgcgtcg	agcctgcccgg	gtttctaccg	caccagcctg	2160
352	accctggccg	cgcggagggc	ggcgggcccc	gtcgaacggc	tgatcgccca	tccgctgcgg	2220
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356	ggctggccgc	tggccgagcg	caccgtggtg	attccctcg	cgatccccac	cgaccggcgc	2340
358	aacgtcgccg	gchgacctcga	cccggtccagc	atccccgaca	aggaacaggc	gatcagcgcc	2400
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364 <211> LENGTH: 1233

364 <211> LENGTH: 1
365 <212> TYPE: DNA

366 <213> ORGANISM: Artificial Sequence

W--> 367 <220> FEATURE:

368 <223> OTHER INFORMATION: pKL4 plasmid coding sequence

368 <225> OTHER -

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375 atgtattggg ttgcgcagac tccagagaag aggctggagt gggtcgcata cattagtaat
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416 <212> TYPE: DNA

417 <213> ORGANISM: Artificial Sequence

W--> 418 <220> FEATURE:

419 <223> OTHER INFORMATION: pMC74 plasmid full sequence

421 <400> SEQUENCE: 7

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426	gcctggagg	gtccctgaaa	ctctcctgt	caacctctgg	attcaacttc	agtgactatt	180
428	acatgtatt	ggttcgccag	actccagaga	agaggctgg	gtgggtcgca	tacatttagta	240
430	atqatgata	ttccggcgct	tattcagaca	ctgtaaagg	ccggttcacc	atctccagag	300

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L:174 M:283 W: Missing Blank Line separator, <220> field identifier
L:243 M:283 W: Missing Blank Line separator, <220> field identifier
L:276 M:283 W: Missing Blank Line separator, <220> field identifier
L:367 M:283 W: Missing Blank Line separator, <220> field identifier
L:418 M:283 W: Missing Blank Line separator, <220> field identifier
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L:937 M:283 W: Missing Blank Line separator, <220> field identifier
L:1070 M:283 W: Missing Blank Line separator, <220> field identifier
L:1265 M:283 W: Missing Blank Line separator, <220> field identifier